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PCT10

RAW SEQUENCE LISTING

PATENT APPLICATION: US/10/088,143

DATE: 08/29/2002

TIME: 14:03:31

Input Set : A:\271\_294.ST25.txt

Output Set: N:\CRF3\08292002\J088143.raw

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3 <110> APPLICANT: PESTKA, Sidney
4   KOTENKO, Sergei
6 <120> TITLE OF INVENTION: Cytomegalovirus-Encoded IL-10 Homolog
8 <130> FILE REFERENCE: 271/294
10 <140> CURRENT APPLICATION NUMBER: 10/088,143
11 <141> CURRENT FILING DATE: 2002-02-22
13 <150> PRIOR APPLICATION NUMBER: PCT/US00/24213
14 <151> PRIOR FILING DATE: 1999-09-02
16 <150> PRIOR APPLICATION NUMBER: 60/152,062
17 <151> PRIOR FILING DATE: 1999-09-02
19 <160> NUMBER OF SEQ ID NOS: 21
21 <170> SOFTWARE: PatentIn version 3.1
23 <210> SEQ ID NO: 1
24 <211> LENGTH: 789
25 <212> TYPE: DNA
26 <213> ORGANISM: Cytomegalovirus
28 <400> SEQUENCE: 1
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31 tctctggtcc tgatcgtctt ttttctaggc gcttccgagg aggcgaagcc ggcgacgacg      120
33 acgataaaga atacaaagcc gcagtgtcgt ccagaggatt acgcgaccag attgcaagat      180
35 ctccgcgtca cctttcatcg agtaaaacct acgttggtag gtcacgtagg tacggtttat      240
37 tgcgacggtc tttcttttcc gcgtgtcggg tgacgtagtt ttctcttctgt agcaacgtga      300
39 ggacgactac tccgtgtggc tcgacggtac ggtggtcaaa ggctgttggg gatgcagcgt      360
41 catggactgg ttgttgaggc ggtatctgga gatcgtgttt cccgcaggcg accacgtcta      420
43 tcccggactc aagacggaat tgcatagtat gcgctcgacg ctagaatcca tctacaaaga      480
45 catgcggaac tgtgtaagtg tctctgtggc ggcgctgtcc gcacagaggt aacaacgtgt      540
47 tcatagcacg ctgttttact tttgtcgggc tcccagcctc tgttaggttg cggagataag      600
49 tccgtgatta gtcggctgtc tcaggaggcg gaaaggaaat cggataacgg cagcggaag      660
51 ggtctcagcg agttggacac gttgttttagc cgtctcgaag agtatctgca ctcgagaaag      720
53 tagcgttgcg atttgacgtc cgctccgggtg tcgttcaccc agttacttta ataaacgtac      780
55 tgtttaacc                                     789
58 <210> SEQ ID NO: 2
59 <211> LENGTH: 175
60 <212> TYPE: PRT
61 <213> ORGANISM: Cytomegalovirus
63 <400> SEQUENCE: 2
65 Met Leu Ser Val Met Val Ser Ser Ser Leu Val Leu Ile Val Phe Phe
66 1                               5                               10                               15
69 Leu Gly Ala Ser Glu Glu Ala Lys Pro Ala Thr Thr Thr Ile Lys Asn
70                               20                               25                               30
73 Thr Lys Pro Gln Cys Arg Pro Glu Asp Tyr Ala Thr Arg Leu Gln Asp
74                               35                               40                               45
77 Leu Arg Val Thr Phe His Arg Val Lys Pro Thr Leu Gln Arg Glu Asp

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78      50      55      60
81 Asp Tyr Ser Val Trp Leu Asp Gly Thr Val Val Lys Gly Cys Trp Gly
82 65      70      75      80
85 Cys Ser Val Met Asp Trp Leu Leu Arg Arg Tyr Leu Glu Ile Val Phe
86      85      90      95
89 Pro Ala Gly Asp His Val Tyr Pro Gly Leu Lys Thr Glu Leu His Ser
90      100      105      110
93 Met Arg Ser Thr Leu Glu Ser Ile Tyr Lys Asp Met Arg Gln Cys Pro
94      115      120      125
97 Leu Leu Gly Cys Gly Asp Lys Ser Val Ile Ser Arg Leu Ser Gln Glu
98      130      135      140
101 Ala Glu Arg Lys Ser Asp Asn Gly Thr Arg Lys Gly Leu Ser Glu Leu
102 145      150      155      160
105 Asp Thr Leu Phe Ser Arg Leu Glu Glu Tyr Leu His Ser Arg Lys
106      165      170      175
109 <210> SEQ ID NO: 3
110 <211> LENGTH: 25
111 <212> TYPE: DNA
112 <213> ORGANISM: Artificial sequence
114 <220> FEATURE:
115 <223> OTHER INFORMATION: synthetic sequence
117 <400> SEQUENCE: 3
118 tcctacagaa actattctaa ccgcg                                25
121 <210> SEQ ID NO: 4
122 <211> LENGTH: 25
123 <212> TYPE: DNA
124 <213> ORGANISM: Artificial sequence
126 <220> FEATURE:
127 <223> OTHER INFORMATION: synthetic sequence
129 <400> SEQUENCE: 4
130 tcattctttcc agcccgcccta gcaac                                25
133 <210> SEQ ID NO: 5
134 <211> LENGTH: 25
135 <212> TYPE: DNA
136 <213> ORGANISM: Artificial sequence
138 <220> FEATURE:
139 <223> OTHER INFORMATION: synthetic sequence
141 <400> SEQUENCE: 5
142 catcataaca taaaggacca cctac                                25
145 <210> SEQ ID NO: 6
146 <211> LENGTH: 25
147 <212> TYPE: DNA
148 <213> ORGANISM: Artificial sequence
150 <220> FEATURE:
151 <223> OTHER INFORMATION: synthetic sequence
153 <400> SEQUENCE: 6
154 cgacgcaacg tggtaaaca gtacg                                25
157 <210> SEQ ID NO: 7
158 <211> LENGTH: 25

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159 <212> TYPE: DNA
160 <213> ORGANISM: Artificial sequence
162 <220> FEATURE:
163 <223> OTHER INFORMATION: synthetic sequence
165 <400> SEQUENCE: 7
166 gggacggcat gctgcggcga tgctg 25
169 <210> SEQ ID NO: 8
170 <211> LENGTH: 25
171 <212> TYPE: DNA
172 <213> ORGANISM: Artificial sequence
174 <220> FEATURE:
175 <223> OTHER INFORMATION: synthetic sequence
177 <400> SEQUENCE: 8
178 agtaactggg tgaacgacac cggag 25
181 <210> SEQ ID NO: 9
182 <211> LENGTH: 25
183 <212> TYPE: DNA
184 <213> ORGANISM: Artificial sequence
186 <220> FEATURE:
187 <223> OTHER INFORMATION: synthetic sequence
189 <400> SEQUENCE: 9
190 aggcgcttcc gaggaggcga agccg 25
193 <210> SEQ ID NO: 10
194 <211> LENGTH: 25
195 <212> TYPE: DNA
196 <213> ORGANISM: Artificial sequence
198 <220> FEATURE:
199 <223> OTHER INFORMATION: synthetic sequence
201 <400> SEQUENCE: 10
202 gactgcaa at cgcaacgcta ctttc 25
205 <210> SEQ ID NO: 11
206 <211> LENGTH: 24
207 <212> TYPE: DNA
208 <213> ORGANISM: Artificial sequence
210 <220> FEATURE:
211 <223> OTHER INFORMATION: synthetic sequence
213 <400> SEQUENCE: 11
214 taggatcctt ccgaggaggc gaag 24
217 <210> SEQ ID NO: 12
218 <211> LENGTH: 20
219 <212> TYPE: DNA
220 <213> ORGANISM: Artificial sequence
222 <220> FEATURE:
223 <223> OTHER INFORMATION: synthetic sequence
225 <400> SEQUENCE: 12
226 atgaattcgt tgttacctct 20
229 <210> SEQ ID NO: 13
230 <211> LENGTH: 22
231 <212> TYPE: DNA

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232 <213> ORGANISM: Artificial sequence
234 <220> FEATURE:
235 <223> OTHER INFORMATION: synthetic sequence
237 <400> SEQUENCE: 13
238 agcggaattc aaatcgcaac gc 22
241 <210> SEQ ID NO: 14
242 <211> LENGTH: 27
243 <212> TYPE: DNA
244 <213> ORGANISM: Artificial sequence
246 <220> FEATURE:
247 <223> OTHER INFORMATION: synthetic sequence
249 <400> SEQUENCE: 14
250 cgggatccca gccagggca gggcacc 27
253 <210> SEQ ID NO: 15
254 <211> LENGTH: 26
255 <212> TYPE: DNA
256 <213> ORGANISM: Artificial sequence
258 <220> FEATURE:
259 <223> OTHER INFORMATION: synthetic sequence
261 <400> SEQUENCE: 15
262 gctctagatc agtttcgtat cttcat 26
265 <210> SEQ ID NO: 16
266 <211> LENGTH: 21
267 <212> TYPE: DNA
268 <213> ORGANISM: Artificial sequence
270 <220> FEATURE:
271 <223> OTHER INFORMATION: synthetic sequence
273 <400> SEQUENCE: 16
274 cgggatcctg cggcgatgct g 21
277 <210> SEQ ID NO: 17
278 <211> LENGTH: 22
279 <212> TYPE: DNA
280 <213> ORGANISM: Artificial sequence
282 <220> FEATURE:
283 <223> OTHER INFORMATION: synthetic sequence
285 <400> SEQUENCE: 17
286 agcggaattc aaatcgcaac gc 22
289 <210> SEQ ID NO: 18
290 <211> LENGTH: 78
291 <212> TYPE: PRT
292 <213> ORGANISM: Cytomegalovirus
294 <400> SEQUENCE: 18
296 Met Leu Ser Val Met Val Ser Ser Ser Leu Val Leu Ile Val Phe Phe
297 1 5 10 15
300 Leu Gly Ala Ser Glu Glu Ala Lys Pro Ala Thr Thr Thr Ile Lys Asn
301 20 25 30
304 Thr Lys Pro Gln Cys Arg Pro Glu Asp Tyr Ala Thr Arg Leu Gln Asp
305 35 40 45
308 Leu Arg Val Thr Phe His Arg Val Lys Pro Thr Leu Val Gly His Val

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309      50                      55                      60
312 Gly Thr Val Tyr Cys Asp Gly Leu Ser Phe Pro Arg Val Gly
313 65                      70                      75
316 <210> SEQ ID NO: 19
317 <211> LENGTH: 178
318 <212> TYPE: PRT
319 <213> ORGANISM: Homo sapiens
321 <400> SEQUENCE: 19
323 Met His Ser Ser Ala Leu Leu Cys Cys Leu Val Leu Leu Thr Gly Val
324 1                      5                      10                      15
327 Arg Ala Ser Pro Gly Gln Gly Thr Gln Ser Glu Asn Ser Cys Thr His
328                      20                      25                      30
331 Phe Pro Gly Asn Leu Pro Asn Met Leu Arg Asp Leu Arg Asp Ala Phe
332                      35                      40                      45
335 Ser Arg Val Lys Thr Phe Phe Gln Met Lys Asp Gln Leu Asp Asn Leu
336      50                      55                      60
339 Leu Leu Lys Glu Ser Leu Leu Glu Asp Phe Lys Gly Tyr Leu Gly Cys
340 65                      70                      75                      80
343 Gln Ala Leu Ser Glu Met Ile Gln Phe Tyr Leu Glu Glu Val Met Pro
344                      85                      90                      95
347 Gln Ala Glu Asn Gln Asp Pro Asp Ile Lys Ala His Val Asn Ser Leu
348                      100                     105                     110
351 Gly Glu Asn Leu Lys Thr Leu Arg Leu Arg Leu Arg Arg Cys His Arg
352                      115                     120                     125
355 Phe Leu Pro Cys Glu Asn Lys Ser Lys Ala Val Glu Gln Val Lys Asn
356      130                     135                     140
359 Ala Phe Asn Lys Leu Gln Glu Lys Gly Ile Tyr Lys Ala Met Ser Glu
360 145                     150                     155                     160
363 Phe Asp Ile Phe Ile Asn Tyr Ile Glu Ala Tyr Met Thr Met Lys Ile
364                      165                     170                     175
367 Arg Asn
371 <210> SEQ ID NO: 20
372 <211> LENGTH: 160
373 <212> TYPE: PRT
374 <213> ORGANISM: Epstein-Barr virus
376 <400> SEQUENCE: 20
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379 1                      5                      10                      15
382 Leu Ala Pro Glu Cys Gly Gly Thr Asp Gln Cys Asp Asn Phe Pro Gln
383                      20                      25                      30
386 Met Leu Arg Asp Leu Arg Asp Ala Phe Ser Arg Val Lys Thr Phe Phe
387                      35                      40                      45
390 Gln Thr Lys Asp Glu Val Asp Asn Leu Leu Leu Lys Glu Ser Leu Leu
391      50                      55                      60
394 Glu Asp Phe Lys Gly Tyr Leu Gly Cys Gln Ala Leu Ser Glu Met Ile
395 65                      70                      75                      80
398 Gln Phe Tyr Leu Glu Glu Val Met Pro Gln Ala Glu Asn Gln Asp Pro
399                      85                      90                      95
402 Glu Ala Lys Asp His Val Asn Ser Leu Gly Glu Asn Leu Lys Thr Leu

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## VERIFICATION SUMMARY

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